

1	MPMPSRDGG LHP RHHH YGSH SPWSQLLSSPMTPTSIKGLYYRRVRKVGAL	hKv6.2
1	MPMSSRDRLHPGH HFGSCSFLSQLWPGE-EPKSVKGLYYSPARKVGNQ	mKv6.2
51	DASEVDLKKEILINVGGRRYLLPWSTLDRFPLSRLSKLRLCRSYEEIVQL	hKv6.2
51	DASFEANLKEILINVGGORYLLPWSTLDAFPLSRLSRLRLCRSHEEITQL	mKv6.2
101	CDDYDEDSQEFFDRSPSAFGVIVSFLAAGKLVLLQEMCALSFOEELAYW	hKv6.2
100	CDDYDEDSQEFFDRNPSTAFGVIVSFLAAGKLVLLREMCALSFOEELSYW	mKv6.2
151	GIEEAHLERCCLRKLLRKLEELLELAKLHREDVLR--QQRERTRRPASHSS	hKv6.2
150	GIEETNLERCCLRKLLKLEEAEE---LRREEAAQROQQRQACHSEVQAS	mKv6.2
199	RWGLCMNRLREMVENPQSGLPQKVFACLSILFVATTAVSLCVSTMPDLRA	hKv6.2
197	RWARSMNQLREMVEDPQSGLPQKVFACLSVLFVATTAVSLCVSTMPDFRA	mKv6.2
249	EEDQGECSRKCYIIFIVETICVAWFSLEFCLRFVQAQDKCQFFQGPLNIIT	hKv6.2
247	EEGKGECTRKCYYIFMVESICVAWFSLEFCLRFVQAPNKCQFFRGPLNVI	mKv6.2
299	DILAISPYVSLAVSEEPPEGERPSRSSLYEKVGLVLRVLRALRILYVM	hKv6.2
297	DILAISPYVSLAVSDESPEAGERPSSSSLYEKVGLVLRVLRALRILYVM	mKv6.2
349	RLARHSLGLQTLGLTVRRC TCEFGLL LFLAVAITLFSPLVYVAEKESGR	hKv6.2
347	RLARHSLGLQTLGLTVRRCAREFGLLMLFLAVAVTLFSPLVYVAENESGR	mKv6.2
399	VLEFTSIPASYWWAIISM TTVGYGDMVPRSVPGQMVALSSILSGILIMAF	hKv6.2
387	VLEFTSIPASYWWAIISM TTVGYGDMVPRSVPGQMVALSSILSGILIMAF	mKv6.2
449	PATSI FHTFSHSYLELKKKEQEQVQARLRRLONTGPASECELLDPHVASEH	hKv6.2
447	PATSI FHTFSHSYLELKKKEQEQVQARLRRLONTNSASE-----R	mKv6.2
499	EIMNDVNDLILEGPALPIMHM	hKv6.2
486	ELLSDVDLVP EGLTSEGRYM	mKv6.2

FIG. 1.

+

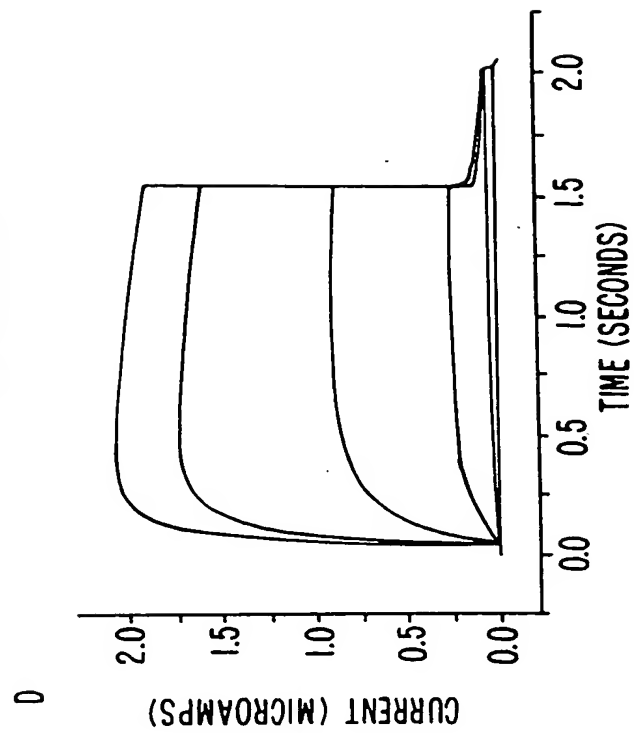
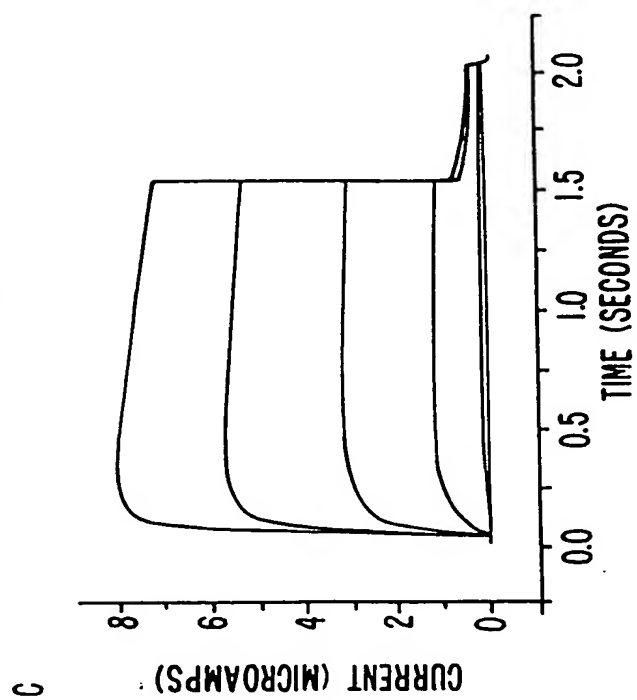
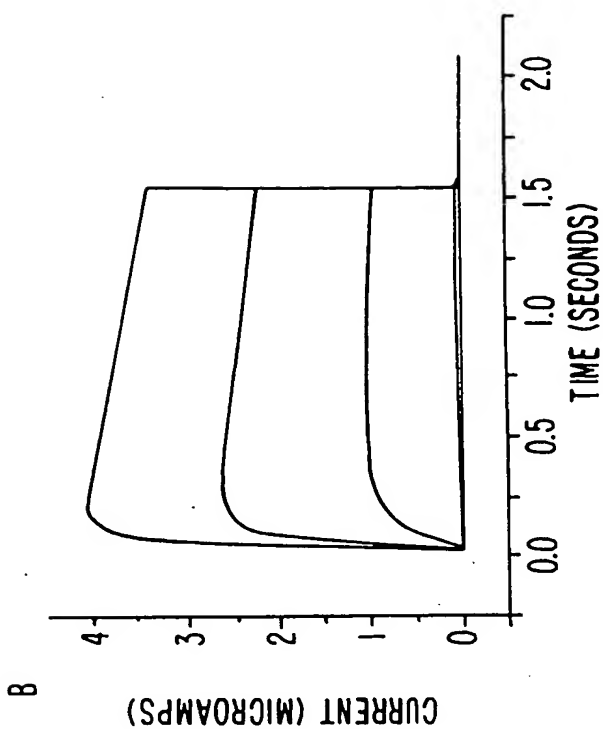
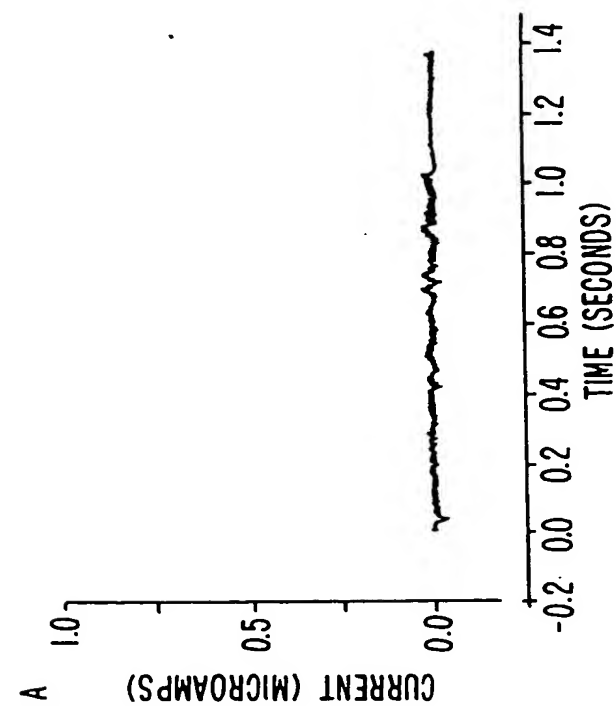


FIG. 2

+